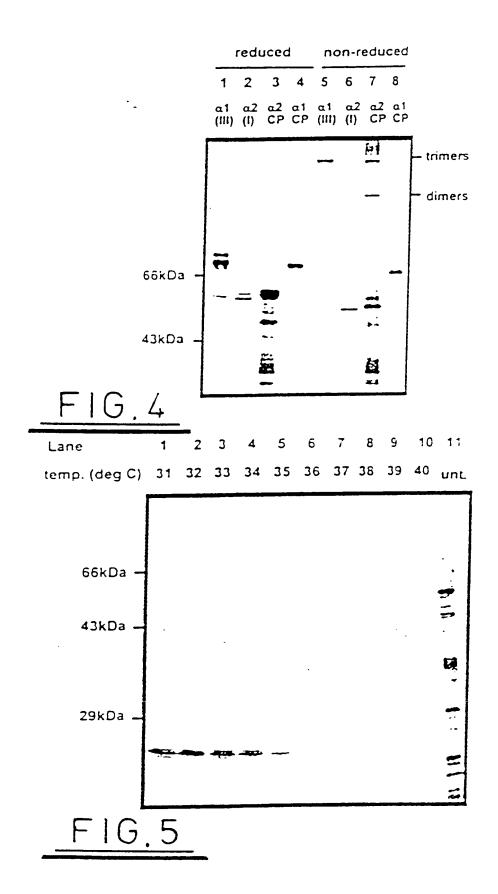
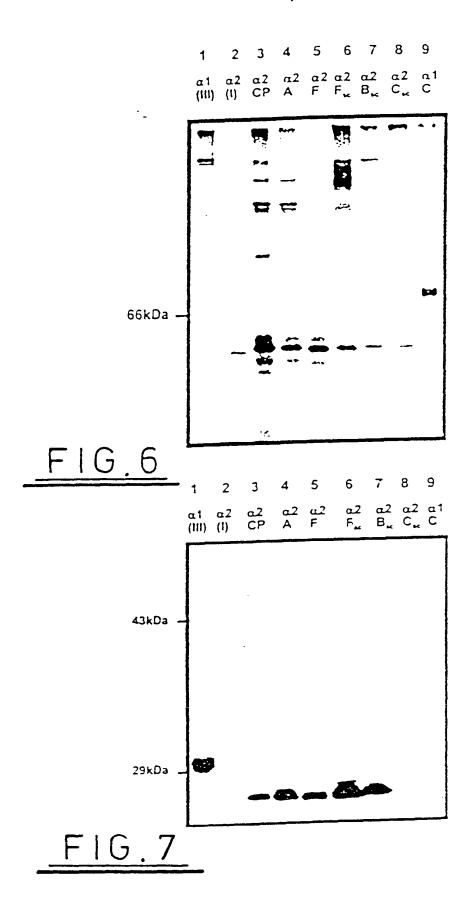
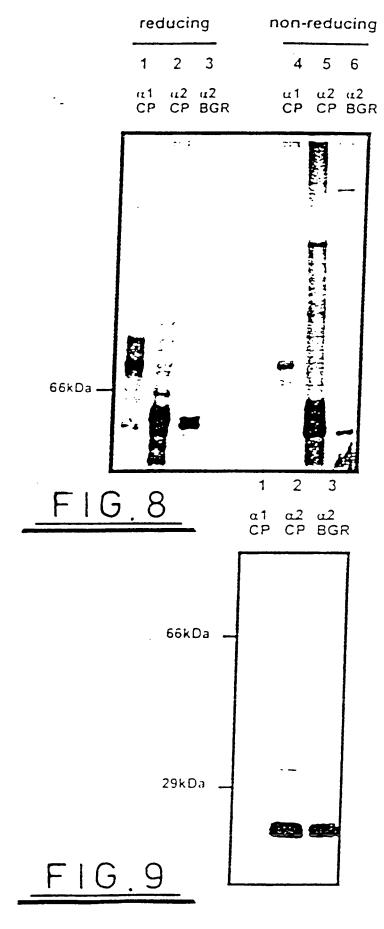


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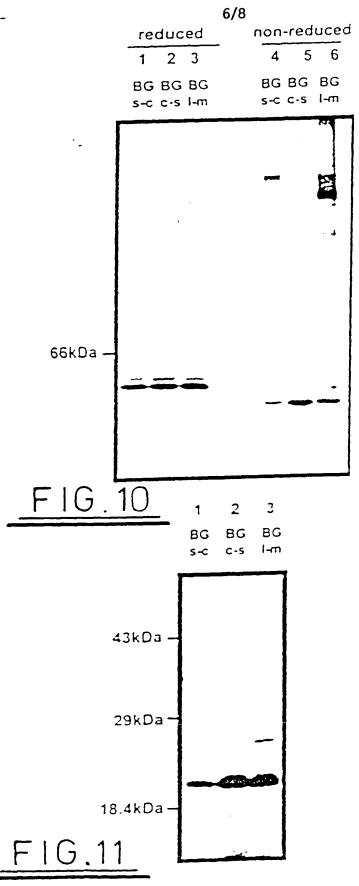
alphal(I) alpha2(I) alphal(III)	YYRADDA NVVRDRDLEV DTTLKSLSQQ IENIRSPEGS RKNPARTCRD LKMCHS FYRADQPRSA PSLRPKDYEV DATLKSLNNQ IETLLTPEGS RKNPARTCRD LRLSHP YYGDEP MDFKINTDEI MTSLKSVNGQ IESLISPDGS RKNPARNCRD LKFCHP	EWSS
alphal(I) alpha2(I) alphal(III)	GEYWIDPNQG CNLDAIKVFC NMETGETCVY PTQPSVAQKN WYISKNPKDK RHVWFG GYYWIDPNQG CTMEAIKVYC DFPTGETCIR AQPENIPAKN WYRSSKDK KHVWLG GEYWVDPNQG CKLDAIKVFC NMETGETCIS ANPLNVPRKH WW.TDSSAEK KHVWFG	ETIN ESMD
alphal(I) alpha2(I) alphal(III)	B C G 6 DGF0FEYGGQ GSDPADVAIQ LTFLRIMSTE ASQNITYHCK NSVAYMDQOT GNLKKAI AGSQFEYNVE GVTSKEMATQ LAFMELLANY ASQNITYHCK NSIAYMDEET GNLKKAV GGFQFSYGNP ELPEDVLDVQ LAFLRLISSR ASQNITYHCK NSIAYMDQAS GNVKKAI	/ILQ
alphal(I) alpha2(I) alphal(III)	GSNEIEIRAE GNSRFTYSVT VDGCTSHTGA WGKTVIEYKT TKTSRLPIID VAPLDVG GSNDVELVAE GNSRFTYTVL VDGGSKKTNE WGKTIIEYKT NKPSRLPFLD IAPLDIG GSNEGEFKAE GNSKFTYTVL EDGCTMHTGE WSKTVFEYRT RKAVRLPIVD IAPYDIG	GAD
alphal(I) alpha2(I) alphal(III)	QEFGFDVGPV CFL HEFFVDIGPV CFK QEFGVDVGPV CFL -111 1-111	
N-propeptide	triple helical domain C-propeptide	
2 360000	proα1(III)Δ1	
	proα2(I)Δ1	
12 Maritistical	proα1(III):(I)CP	
	proα2(I):(III)CP	
	proα2(I):(III)A	
	proα2(I):(III)F	
	proα2(I):(III)B	
	proα2(I):(III)C	
2 carisconnyduaen	proα1(III):(I)C	
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<u> </u>	FIG. 3	

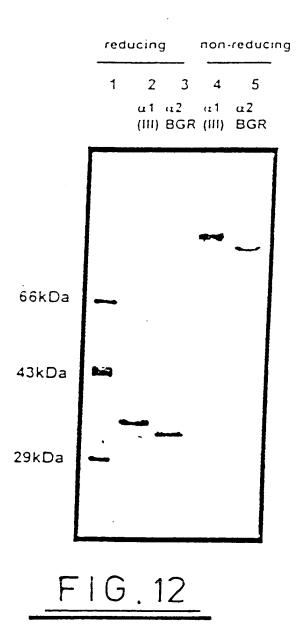






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	alphal(I)	alpha2(I)	alpha1(II)	alphal(III)	alphal(V)	alpha2 (V)	alphal(XI)	alpha2(XI)

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